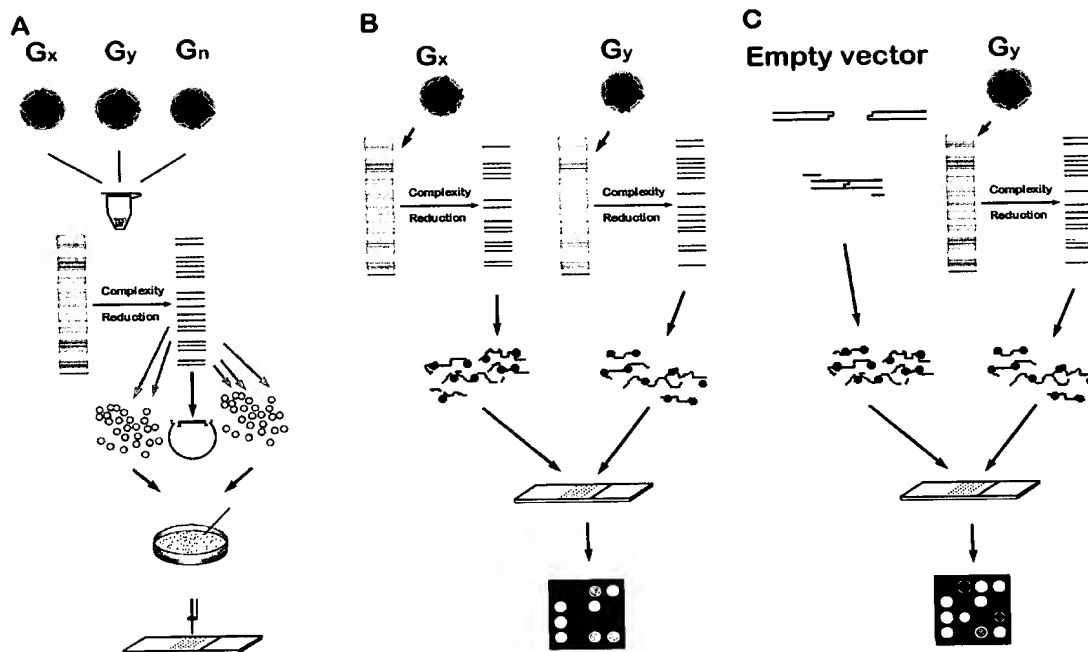
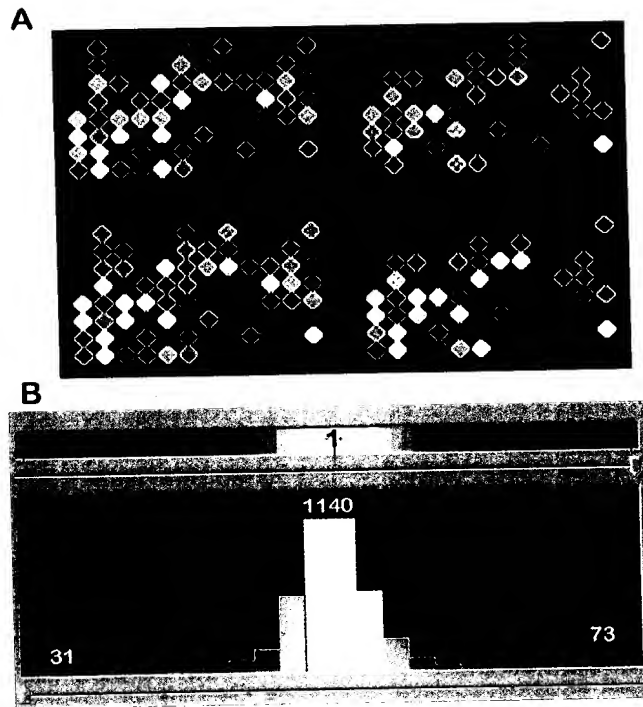


FIGURE 1



**FIGURE 2**



**FIGURE 3**

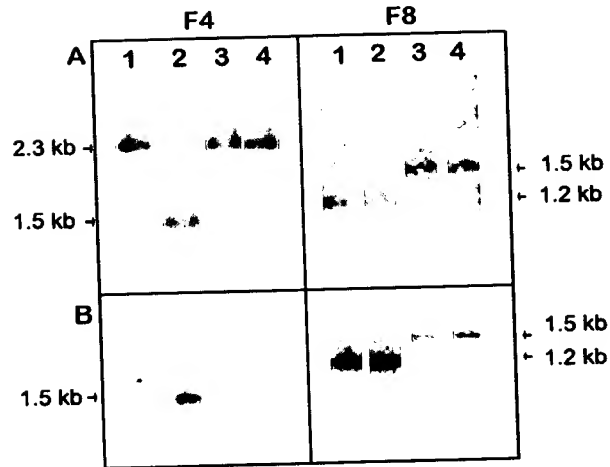




Figure 4

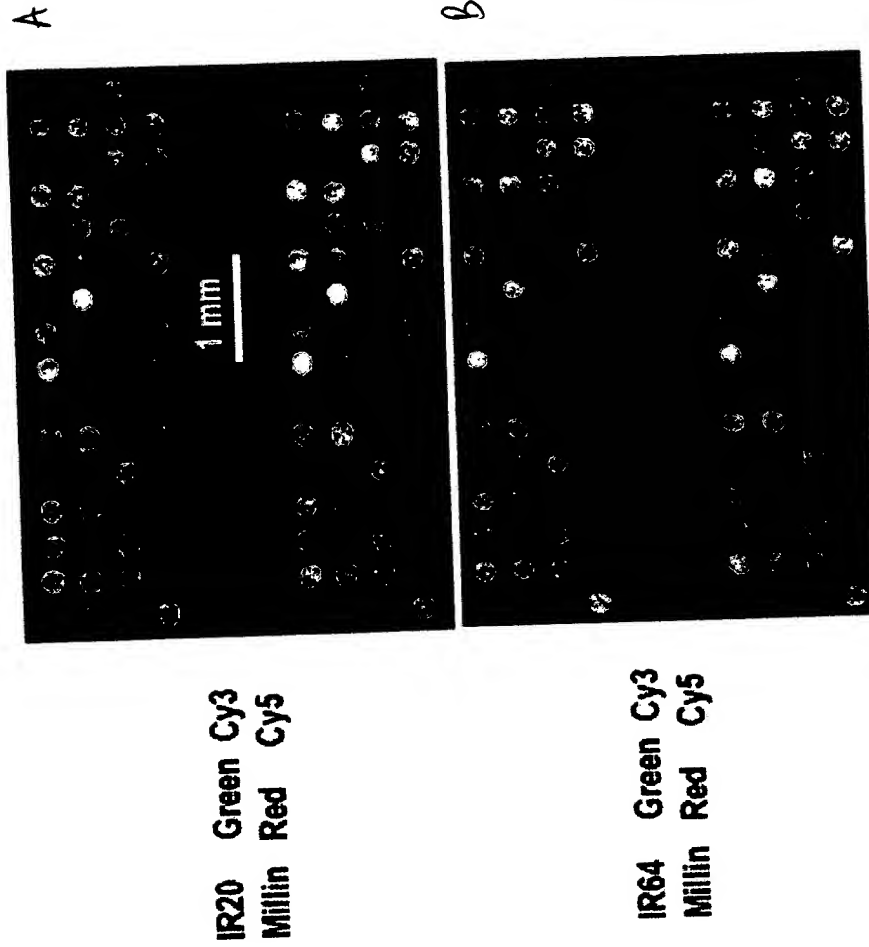
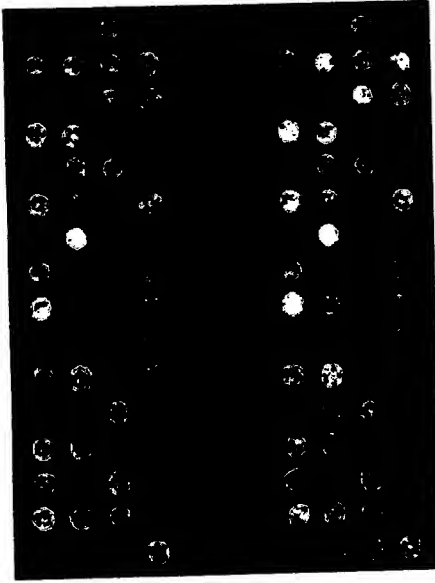


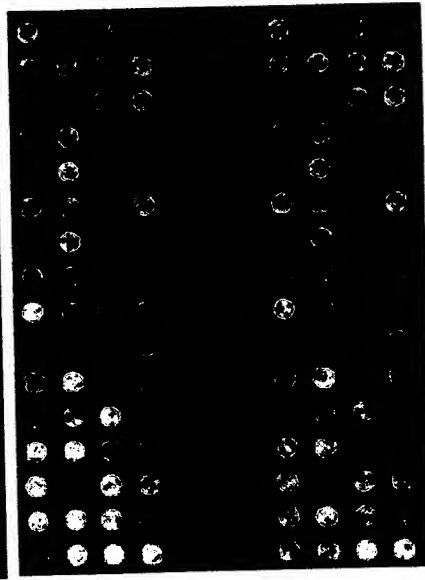
FIGURE 5

A



IR20 Green Cy3  
Millin Red Cy5

B

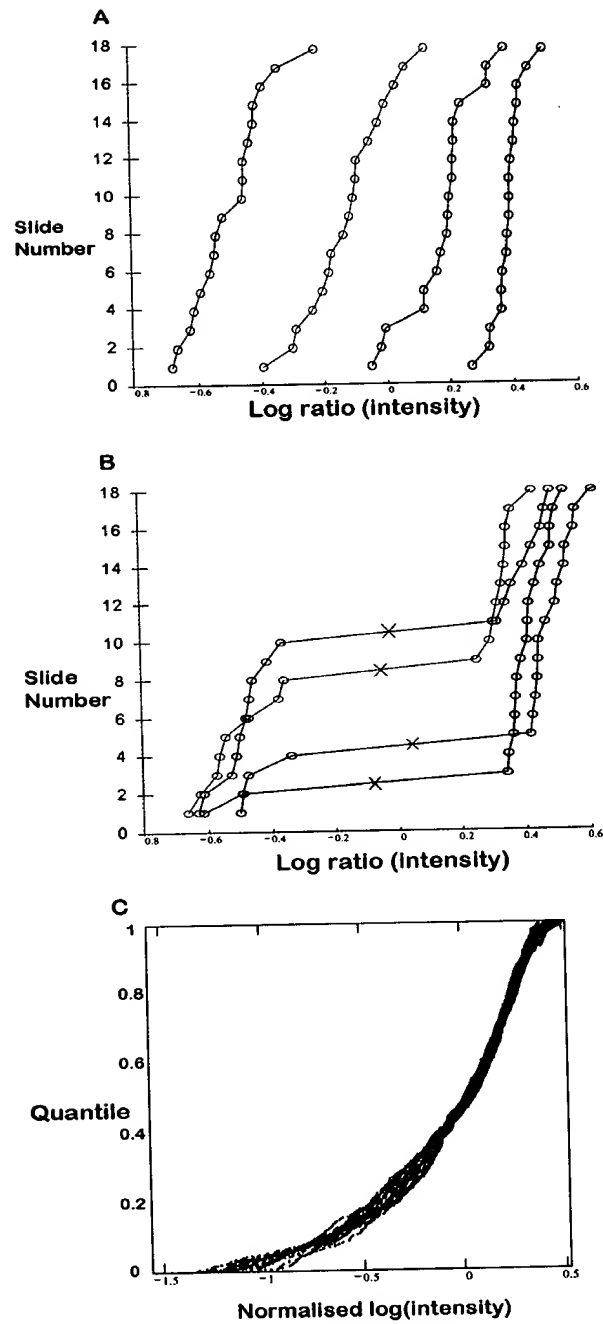


IR20 Green Cy3  
Vector Red Cy5

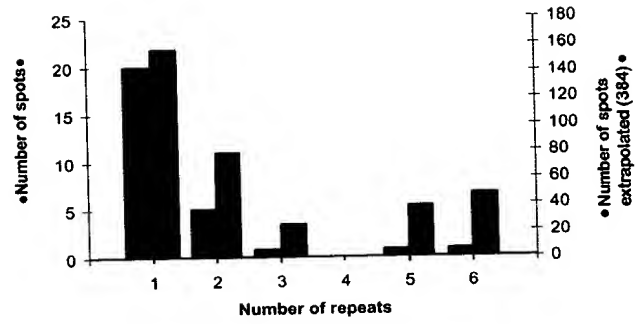
1 mm

FIGURE 6

FIGURE 7

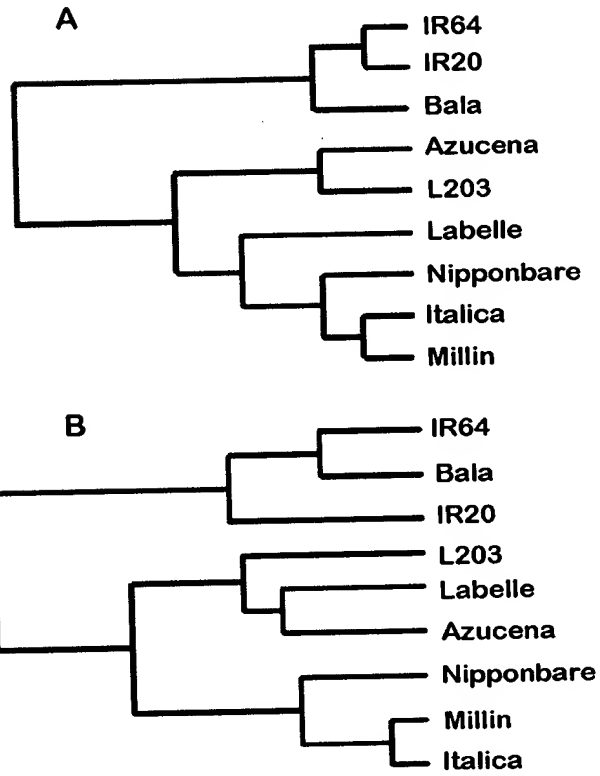


**FIGURE 8**

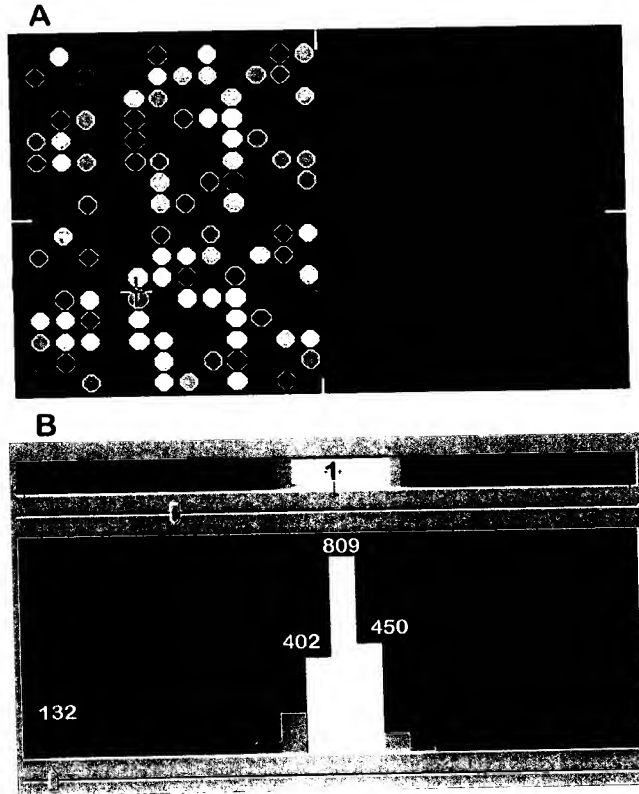




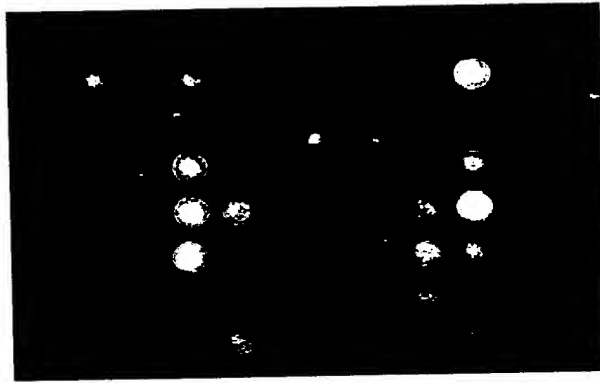
**FIGURE 9**



**FIGURE 10**



## Barley DArT polymorphism analysis between Steptoe and Morex cultivars



Morex Diversity Panel was labelled with Cy3 and Steptoe Diversity Panel with Cy5 dyes

## Mouse DArTs using cDNA-based Diversity Array

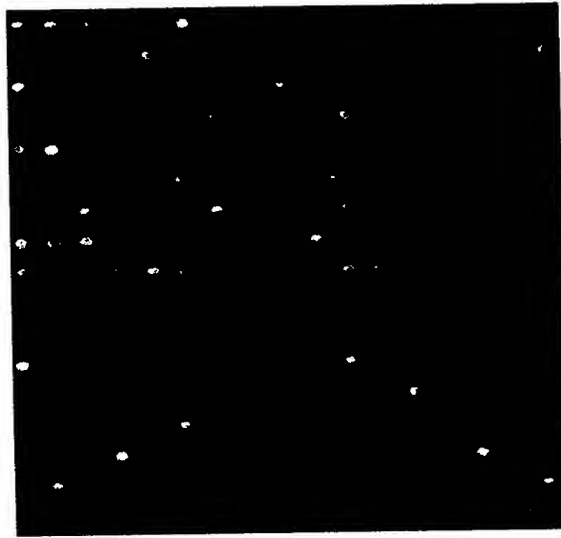
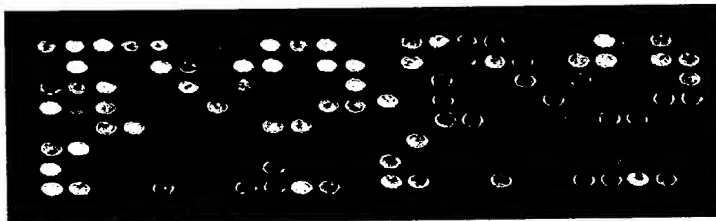


Figure 12

Methylation pattern differences among rice tissues typed using  
DArTs



Red: callus  
Green: seedling root  
Yellow: callus + root



Red: callus  
Green: immature embryo  
Yellow: callus + embryo

## Amplicon Southern analysis

1 2 3 4 5 6 7 8 9 10 11 12



### Rice Tissue

### Log Intensity(Diversity Panel/TOPO)

1. Seedling leaves	0.51
2. Seedling root	0.64
3. Mature pollen and anther	0.58
4. immature pollen and anther	0.34
5. Fertilized ovary and stigma	-0.01
6. Unfertilized ovary and stigma	0.39
7. Mature embryo	0.37
8. Immature embryo	0.47
9. Immature endosperm	0.42
10. Flag leaves	0.51
11. Callus	0.60
12. DNA Mixture	0.63